

AJ

SEQUENCE LISTING



<110> Burch, Ronald
Sackler, David

<120> Contraceptive Antibody Vaccines

<130> 6750-018-999

<140> 09/831,631

<141> 2001-05-10

<160> 70

<170> PatentIn version 3.0

<210> 1

<211> 16

<212> DNA

<213> Artificial Sequence

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<223> Description of Artificial Sequence: Primer for PCR

<400> 1
aacagctatg accatg 16

<210> 2

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

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<223> Description of Artificial Sequence: Primer for PCR

<400> 2
gaattcatgg cttgggtgtg 20

<210> 3

<211> 13

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<223> Description of Artificial Sequence: CDR Derived peptide with Biotin label at the N-terminal residue

<400> 3
Thr Ala Lys Ala Ser Gln Ser Val Ser Asn Asp Val Ala
1 5 10

<210> 4

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

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<223> Description of Artificial Sequence: CDR Derived peptide with Biotin

label at the N-terminal residue

<400> 4

Ile Tyr Tyr Ala Ser Asn Arg Tyr Thr
1 5

<210> 5

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<221> misc_feature

<223> Description of Artificial Sequence: CDR Derived peptide with Biotin
label at the N-terminal residue

<400> 5

Phe Ala Gln Gln Asp Tyr Ser Ser Pro Leu Thr
1 5 10

<210> 6

<211> 7

<212> PRT

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<223> Description of Artificial Sequence: CDR Derived peptide with Biotin
label at the N-terminal residue

<400> 6

Phe Thr Asn Tyr Gly Met Asn
1 5

<210> 7

<211> 19

<212> PRT

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<220>

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<223> Description of Artificial Sequence: CDR Derived peptide with Biotin
label at the N-terminal residue

<400> 7

Ala Gly Trp Ile Asn Thr Tyr Thr Gly Glu Pro Thr Tyr Ala Asp
1 5 10 15
Asp Phe Lys Gly

<210> 8

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<221> misc_feature

<223> Description of Artificial Sequence: CDR Derived peptide with Biotin
label at the N-terminal residue

<400> 8

Ala Arg Ala Tyr Tyr Gly Lys Tyr Phe Asp Tyr
1 5 10

<210> 9
<211> 221
<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Sperm cell specific epitope

<400> 9
gaattccagc cttcaggtga acatggctcc ggtgaacagc cttctggtga gcaggcctcg 60
ggtgaacagc cttcaggtga gcacgcttca ggggaacagg cttcaggtgc accaatttca 120
agcacatcta caggcacaat attaaattgc tacacatgtg cttatatgaa tgatcaagga 180
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<210> 10
<211> 69
<212> PRT
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description of Artificial Sequence: Sperm cell specific epitope

<400> 10
Gln Pro Ser Gly Glu His Gly Glu Gln Pro Ser Gly Glu Gln Ala Ser
1 5 10 15
Gly Glu Gln Pro Ser Gly Glu His Ala Ser Gly Glu Gln Ala Ser Gly
20 25 30
Ala Gln Ile Ser Ser Thr Ser Thr Gly Thr Ile Leu Asn Cys Tyr Thr
35 40 45
Cys Ala Tyr Met Asn Asp Gln Gly Lys Cys Leu Arg Gly Glu Gly Thr
50 55 60
Cys Ile Thr Gln Asn
65

<210> 11
<211> 75
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description of Artificial Sequence: Cloning primers for SP10

<400> 11
gaattccagc cttcaggtga acatggctcc ggtgaacagc cttctggtga gcaggcctcg 60
ggtgaacagc cttag 75

<210> 12
<211> 75
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description of Artificial Sequence: Cloning primers for SP10

<400> 12
gtgagcagcgttccaggggaa cagccttcag gtgcaccaat ttcaaggaca tctacaggca 60
caatattaaa ttgct 75

<210> 13
 <211> 70
 <212> DNA
 <213> Artificial Sequence

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 <223> Description of Artificial Sequence: Cloning primers for SP10

 <400> 13
 acacatgtgc ttatatgaat gatcaaggaa aatgtcttcg tggagaggga acctgcatca 60
 ctcagaattc 70

 <210> 14
 <211> 70
 <212> DNA
 <213> Artificial Sequence

 <220>
 <221> misc_feature
 <223> Description of Artificial Sequence: Cloning primers for SP10

 <400> 14
 acacagcagc ttatatgaat gatcaaggaa aagcacttcg tggagaggga accgcaatca 60
 ctcagaattc 70

 <210> 15
 <211> 79
 <212> DNA
 <213> Artificial Sequence

 <220>
 <221> misc_feature
 <223> Description of Artificial Sequence: Cloning primers for SP10

 <400> 15
 gaattctgag tgatgcaggt tccctctcca cgaagacatt ttccttgatc attcatataa 60
 gcacatgtgt agcaattta 79

 <210> 16
 <211> 79
 <212> DNA
 <213> Artificial Sequence

 <220>
 <221> misc_feature
 <223> Description of Artificial Sequence: Cloning primers for SP10

 <400> 16
 gaattctgag tgattgcggt tccctctcca cgaagtgctt tttgatgatc attcatataa 60
 gctgctgtgt agcaattta 79

 <210> 17
 <211> 75
 <212> DNA
 <213> Artificial Sequence

 <220>
 <221> misc_feature
 <223> Description of Artificial Sequence: Cloning primers for SP10

 <400> 17

atattgtgcc ttagatgtg cttgaaattg gtgcacctga agcctgttcc cctgaagcgt 60
gctcacctga aggct 75

<210> 18
<211> 67
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description of Artificial Sequence: Cloning primers for SP10

<400> 18
gttctcccga ggctgtctca ccagaaggct gttcacccga gccatgttca cctgaaggct 60
ggaattc 67

<210> 19
<211> 210
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description of Artificial Sequence: Sperm cell specific epitope M
SA-6

<400> 19
gtcggcagcc tccgaagcag cccgctccag agcccgtgc tccgaccgct cgtccagagc 60
agcctctgct tgctgttct cttgctgcga tacagctgcg gcgacggcag ctgcagccga 120
cgatactgcg acttgacggt gtgccggcga atgtacttgc tgctgcgatt cacggaccg 180
ccgctcccgc agacgtgctg cgtcttgagc 210

<210> 20
<211> 70
<212> PRT
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description of Artificial Sequence: Sperm cell specific epitope M
SA-6

<400> 20
Gln Pro Ser Glu Ala Ser Ser Gly Glu Val Ser Gly Asp Glu Ala Gly
1 5 10 15
Glu Gln Val Ser Ser Glu Thr Asn Asp Lys Glu Asn Asp Ala Met Ser
20 25 30
Thr Pro Leu Pro Ser Thr Ser Ala Ala Ile Thr Leu Asn Cys His Thr
35 40 45
Cys Ala Tyr Met Asn Asp Asp Ala Lys Cys Leu Arg Gly Glu Gly Val
50 55 60
Cys Thr Thr Gln Asn Ser
65 70

<210> 21
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Oligomer from MSA 63

<400> 21
gtcggcagcc tccgaagcag cccgctccag agcccgtgc tccga 45

<210> 22
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description of Artificial Sequence: Oligomer from MSA 63

<400> 22
agcccgtgc tccgaccgct cgtccagagc agcctctgct tgctg 45

<210> 23
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description of Artificial Sequence: Oligomer from MSA 63

<400> 23
agcctctgct tgctgttctt cttgctgcga tacagctgcg gcgac 45

<210> 24
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description of Artificial Sequence: Oligomer from MSA 63

<400> 24
tacagctgcg gcgacggcag ctgcagccga cgatactgcg acttg 45

<210> 25
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Oligomer from MSA 63

<400> 25
cgatactgcg acttgacggt gtgcacgcga atgtacttgc tgctg 45

<210> 26
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Oligomer from MSA 63

<400> 26
atgtacttgc tgctgcgatt cacggacgcg ccgctcccgc agacg 45

<210> 27
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Oligomer from MSA 63

<400> 27
cgattcacgg acgcgccgct cccgcagacg tgctgcgtct tgagc 45

<210> 28
<211> 17
<212> PRT
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Consensus sequence

<400> 28
Gln Pro Ser Glu Ala Ser Ser Gly Glu Val Ser Gly Asp Glu Ala Gly
1 5 10 15
Glu

<210> 29
<211> 384
<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Consensus sequence

<400> 29
atggcttggt gttggacctt gctattcctg atggcagctg cccaaagtgc ccaagcagat 60
atgcaaataa cacaagtcc tagtagtttg agtgctagtg tgggagatca agtgacaatc 120
acatgtcggg ctagtcaaag tatcagtaac tgtttggtt ggtatcaaca aaagcctgga 180
aaggctccta agttgttgat ctatgctgct agtagtttg agagtggagt gcctagtcgg 240
ttcagtggaa gtggaagtgg aacacggttc accttgacca tgagtagttt gcaacctgag 300
gatttcgcta cctattattg tcaacaatat aacagtttgc cttggacctt cggacaagga 360
accaaggtgg agatcaagga attc 384

<210> 30
<211> 417
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description of Artificial Sequence: Consensus sequence

<400> 30
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gttcagctgg tgcagtgtgg cgctgaggtg aagaagcctg gcgcttctgt gaaggtgtct 120
tgcaaggctt ctggctacac attcacatct tacgctatat cttggaattg ggtgaggcag 180

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<210> 31
 <211> 63
 <212> DNA
 <213> Artificial Sequence

<220>
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 <223> Description of Artificial Sequence: Consensus sequence

<400> 31	
gacattgtga	60
tgacacagtc	
tccatcctcc	
ctagctgtgt	
cagttggaga	
gaaggttact	63
atg	

<210> 32
 <211> 74
 <212> DNA
 <213> Artificial Sequence

<220>
 <221> misc_feature
 <223> Description of Artificial Sequence: Consensus sequence

<400> 32	
gcaagctcat	60
agtaaccttc	
tctccaactg	
acacacgata	
gggaggatgg	
agactgtgac	74
atcacaatgt	
ctgc	

<210> 33
 <211> 84
 <212> DNA
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 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

<400> 33	
agctgcgtcg	60
gcagcctccg	
aagcagcccg	
ctccagagcc	
cgctgctggc	
atggtaccag	84
cagaaaccag	
ggcagctctc	
taaa	

<210> 34
 <211> 72
 <212> DNA
 <213> Artificial Sequence

<220>
 <221> misc_feature
 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

<400> 34	
ctgccctggt	60
ttctgctggt	
accatcggag	
cagcgggctc	
tgccggagcgg	
gctgcttcgg	72
acggctgccg	
ac	

<210> 35
 <211> 78

<212> DNA
 <213> Artificial Sequence

 <220>
 <221> misc_feature
 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

 <400> 35
 gacattgtga tgtcacagtc tccatcctcc ctagctgtgt cagttggaga gaagggttact 60
 gtgagcgcta agtccagt 78

 <210> 36
 <211> 75
 <212> DNA
 <213> Artificial Sequence

 <220>
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 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

 <400> 36
 gagagccttt tatatagtag caatcaaaag atctacttgg cctggtacca gcagaaacca 60
 gggcagtctc ctaaa 75

 <210> 37
 <211> 67
 <212> DNA
 <213> Artificial Sequence

 <220>
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 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

 <400> 37
 ctgctgattt actgggcatc cactagggaa tctgggggtcc ctgatcgctt cacaggctgg 60
 atctggg 67

 <210> 38
 <211> 68
 <212> DNA
 <213> Artificial Sequence

 <220>
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 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

 <400> 38
 gcacagcaat attatagata tctcggacgt tcggtggagc caccaagctg caaatcaaac 60
 cggaattc 68

 <210> 39
 <211> 69
 <212> DNA
 <213> Artificial Sequence

 <220>
 <221> misc_feature

<223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

<400> 39
accgcctgtg aagcgatcag gcaccccaaga ttccctagtg gatgcccgat aaatcagcag 60
ttaggaga 69

<210> 40
<211> 77
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
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<400> 40
ctgccctggg ttctgctggg accaggccaa gtagatcttt tgagattgct actatataaa 60
aggctctgac tggactt 77

<210> 41
<211> 78
<212> DNA
<213> Artificial Sequence

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<223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

<400> 41
agcgctcata gtaaccttct ctccaactga cacagctagc gacgatcgag actgtgacat 60
cacaatgtct gcttgggc 78

<210> 42
<211> 78
<212> DNA
<213> Artificial Sequence

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<223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

<400> 42
gaattcccggt ttgatttcca gcttggtgcc tccaccgaac gtccgaggat atctataata 60
ttgctgtgcg taataaac 78

<210> 43
<211> 57
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

<400> 43
 agagatttga gtctgaccat cagcagtgtg aaggctgaag acgtggcagt ttattac 57

 <210> 44
 <211> 57
 <212> DNA
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 <220>
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 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

 <400> 44
 tgccagggtct tcagccttga cactgctgat ggtgagagtg aaatctgtcc cagatcc 57

 <210> 45
 <211> 66
 <212> DNA
 <213> Artificial Sequence

 <220>
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 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

 <400> 45
 tcgtgccagt tcctcgtcga ctagctcttc gactagctcc tgctgctctt gtcggtcacg 60
 gaattc 66

 <210> 46
 <211> 75
 <212> DNA
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 <223> Description for Artificial Sequence: Construct for MSA1 and MSALVAC-1

 <400> 46
 gaattccgtg accgacaaga gcagcaggag ctagtcgaag agctggtcga cgaggaactg 60
 gcacgacggg ttcgt 75

 <210> 47
 <211> 80
 <212> DNA
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 <223> Description for Artificial Sequence: Constructs for LDH-C4

 <400> 47
 gaattcatgg cttgggtgtg gaccttgcta ttcctgatgg cagctgccca aagtgcccaa 60
 gcacagatcc agttgggtgca 80

 <210> 48
 <211> 79

<212> DNA
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 <223> Description for Artificial Sequence: Constructs for LDH-C4

 <400> 48
 gtctggacct gagctgaaga agcctggaga gacagtcaag atctccgcta aggcttctgg 60
 gtataccttc acaaactag 79

 <210> 49
 <211> 80
 <212> DNA
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 <223> Description for Artificial Sequence: Constructs for 2CAVHCOL1

 <400> 49
 gaatgaactg ggtgaagcag gctccaggaa agggttttaa gtggatgggc tggataaaca 60
 cctacactgg agagccaaca 80

 <210> 50
 <211> 80
 <212> DNA
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 <400> 50
 tatgctgatg acttcaaggg acggtttgcc ttctctttgg aaacctctgc cagcactgcc 60
 tatttgcaag atcaacacct 80

 <210> 51
 <211> 70
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 <400> 51
 caaaaatgag gacacggcta catatttcgc tgcaagagcc tactatggta aatactttga 60
 ctacgaattc 70

 <210> 52
 <211> 49
 <212> DNA
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 <223> Description for Artificial Sequence: Constructs for 2CAVHCOL1

 <400> 52
 gaattcgtag tcaaagtatt taccatagta ggctcttgca gcaaatatg 49

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<210> 53
<211> 81
<212> DNA
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<223> Description for Artificial Sequence: Constructs for 2CAVHCOL1

<400> 53
tagcctgtgt ctcatttttt gaggttggtg atctgcaa at aggcagtgtt ggcagaggtt 60
tccaaagaga aggcaaaccg t 81

<210> 54
<211> 80
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
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<400> 54
cccttgaagt catcagcata tggtggctct ccagtgtagg tgtttatcca gcccatccac 60
tttaaaccct ttcctggagc 80

<210> 55
<211> 81
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description for Artificial Sequence: Constructs for 2CAVHCOL1

<400> 55
ctgcttcacc cagttcattc catagtttgt gaagggtatac ccagaagcct tagcggagat 60
cttgactgtc tctccaaggc t 81

<210> 56
<211> 100
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description for Artificial Sequence: Constructs for 2CAVHCOL1

<400> 56
tcttcagctc aggtccagac tgcaccaact ggatctgtgc ttgggcaatt tcggcagctg 60
ccatcaggaa tagcaaggtc cacaccaag ccatgaattc 100

<210> 57
<211> 63
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description for Artificial Sequence: Constructs for 2CAVHCOL1

<400> 57

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agtattgtga tgacccagac tcccaaattc ctgcttgtat cagcaggaga cagggttacc 60
ata 63

<210> 58
<211> 64
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description for Artificial Sequence: Constructs for 2CAVHCOL1

<400> 58
acctgcaagg ccagtcagag tgtgagtaat gatgtagctt ggtaccaaca gaaaaccagg 60
gcag 64

<210> 59
<211> 69
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 59
tctcctaaac tgctgatata ctatgcatcc aatcgctaca ctggagtccc tgatcgcttc 60
actggcagt 69

<210> 60
<211> 64
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 60
ggatatggga cggatttcac tttcaccatc agcactgtgc aaggctgaag acctggcagt 60
ttat 64

<210> 61
<211> 69
<212> DNA
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<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 61
ttctgycagc aggattatag ctctccgctc accttcggtg ctgggaccaa gctggacctg 60
aaagaattc 69

<210> 62
<211> 78
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature

<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 62
gaattctttc agctccagct tgggtcccagc accgaacgtg agcggagagc tataatcctg 60
ctgacagaaa taaactgc 78

<210> 63
<211> 63
<212> DNA
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<400> 63
caggtcttca gcctgcacag tgctgatggt gaaagtgaaa tccgtcccat atccactgcc 60
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<210> 64
<211> 69
<212> DNA
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<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 64
gaagcgatca gggactccag tgtagcgatt ggatgcatag tatatcagca gtttaggaga 60
ctgccctgg 69

<210> 65
<211> 63
<212> DNA
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 65
tttctgttgg taccaagcta catcattact cacactctga ctggccttgc tggttatggt 60
aac 63

<210> 66
<211> 63
<212> DNA
<213> Artificial Sequence

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<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 66
cctgtctcct gctcatacaa gcaggaattt gggagtctgg gtcatacaaa tacttgcttg 60
ggc 63

<210> 67
<211> 68
<212> DNA
<213> Artificial Sequence

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<220>
<221> misc_feature
<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 67
ttcgctcagc aggattatag ctctccgctc acgttcggtg ctgggaccaa gctggagctg      60
aaagaatc                                         68

<210> 68
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<212> DNA
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<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 68
gaattctttc agctccagct tgggtcccagc accgaacgtg agcggagagc tataatcctg      60
ctgagcgaaa taaactgc                                         78

<210> 69
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<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<220>
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<400> 69
atg gct tgg gtg tgg acc ttg cta ttc ctg atg gca gct gcc caa agt      48
Met Ala Trp Val Trp Thr Leu Leu Phe Leu Met Ala Ala Ala Gln Ser
1      5      10      15
gcc caa gca gac att gtg atg tca cag tct cca tcc tcc cta gct gtg      96
Ala Gln Ala Asp Ile Val Met Ser Gln Ser Pro Ser Ser Leu Ala Val
20      25      30

tca gtt gga gag aag gtt act atg agc tgc aag tcc agt cag agc ctt      144
Ser Val Gly Glu Lys Val Thr Met Ser Cys Lys Ser Ser Gln Ser Leu
35      40      45

tta tat agt agc aat caa aag atc tac ttg gcc tgg tac cag cag aaa      192
Leu Tyr Ser Ser Asn Gln Lys Ile Tyr Leu Ala Trp Tyr Gln Gln Lys
50      55      60

cca ggg cag tct cct aaa ctg ctg att tac tgg gca tcc act agg gaa      240
Pro Gly Gln Ser Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu
65      70      75      80

tct ggg gtc cct gat cgc ttc aca ggc ggt gga tct ggg aca gat ttc      288
Ser Gly Val Pro Asp Arg Phe Thr Gly Gly Gly Ser Gly Thr Asp Phe
85      90      95

act ctc acc atc agc agt gtg aag gct gaa gac ctg gca gtt tat tac      336
Thr Leu Thr Ile Ser Ser Val Lys Ala Glu Asp Leu Ala Val Tyr Tyr
100     105     110

tgt cag caa tat tat aga tat cct cgg acg ttc ggt gga ggc acc aag      384
Cys-Gln-Gln-Tyr-Tyr-Arg-Tyr-Pro-Arg-Thr-Phe-Gly-Gly-Gly-Thr-Lys
115     120     125

ctg gaa atc aaa cgg                                         399

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Leu Glu Ile Lys Arg
130

<210> 70
<211> 133
<212> PRT
<213> Artificial Sequence

<220>
<221> misc_feature
<223> Description for Artificial Sequence: Constructs for 2CAVLCOL1

<400> 70
Met Ala Trp Val Trp Thr Leu Leu Phe Leu Met Ala Ala Ala Gln Ser
1 5 10 15
Ala Gln Ala Asp Ile Val Met Ser Gln Ser Pro Ser Ser Leu Ala Val
20 25 30
Ser Val Gly Glu Lys Val Thr Met Ser Cys Lys Ser Ser Gln Ser Leu
35 40 45
Leu Tyr Ser Ser Asn Gln Lys Ile Tyr Leu Ala Trp Tyr Gln Gln Lys
50 55 60
Pro Gly Gln Ser Pro Lys Leu Leu Ile Tyr Trp Ala Ser Thr Arg Glu
65 70 75 80
Ser Gly Val Pro Asp Arg Phe Thr Gly Gly Gly Ser Gly Thr Asp Phe
85 90 95
Thr Leu Thr Ile Ser Ser Val Lys Ala Glu Asp Leu Ala Val Tyr Tyr
100 105 110
Cys Gln Gln Tyr Tyr Arg Tyr Pro Arg Thr Phe Gly Gly Gly Thr Lys
115 120 125
Leu Glu Ile Lys Arg
130